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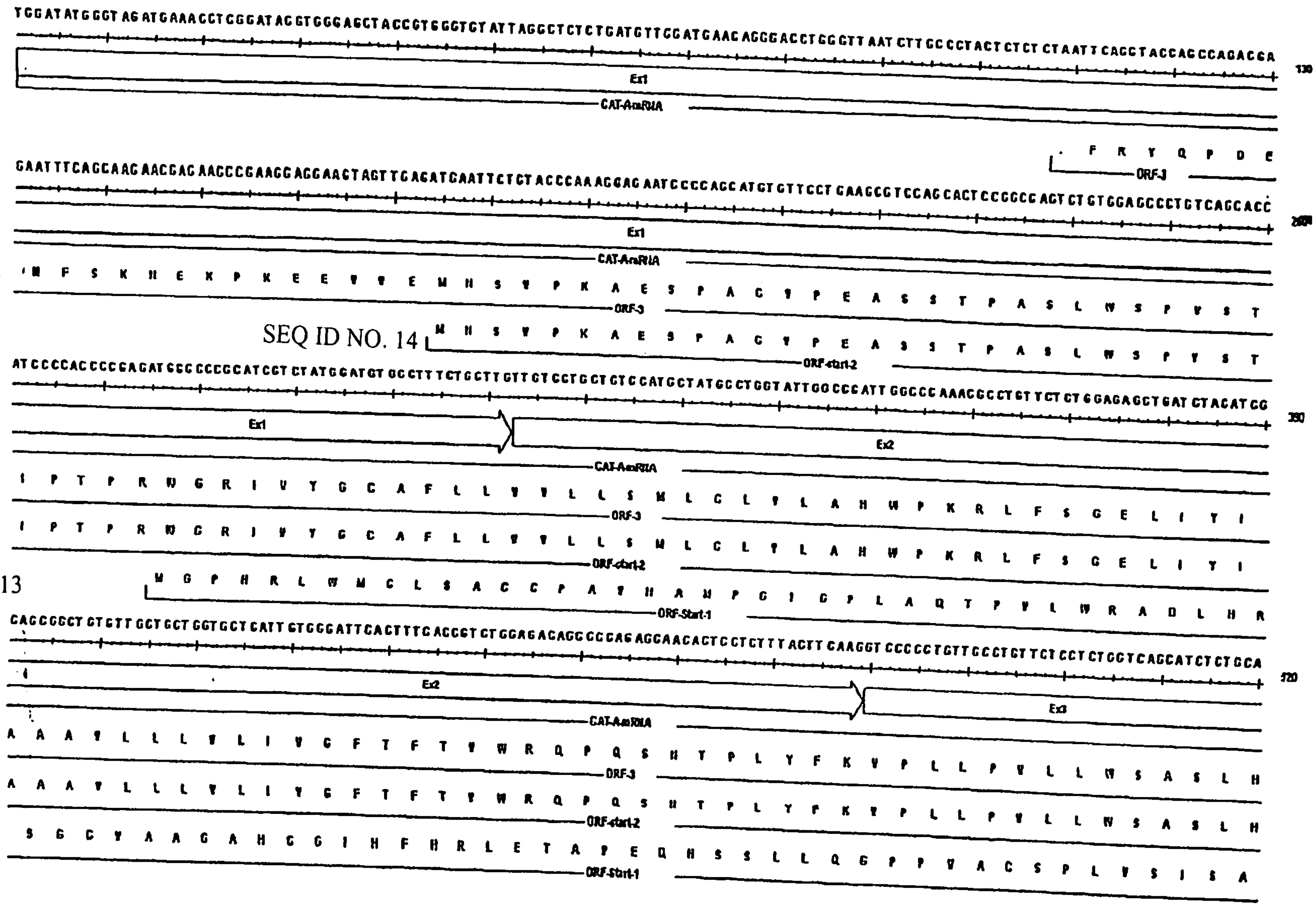
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130
Ex1
CAT-AmRIA
· F R Y Q P D E
ORF-3
GAATTTCAGCAAGAACGAGAACCCGAAGGAGGAACTAGTTGAGATGAATTCTCTACCCAAAGGAGAATCCCCAGCATGTGTTCCTGAAGCGTCCAGCACTCCCGCCAGTCTGTGGAGCCCTGTCAGCACC
260
Ex1
CAT-AmRIA
SEQ ID NO. 15
M F S K N E K P K E E V V E M N S V P K A E S P A C V P E A S S T P A S L W S P V S T
ORF-3
SEQ ID NO. 14
M N S V P K A E S P A C V P E A S S T P A S L W S P V S T
ORF-start-2
ATCCCCACCCCCAGATGGGGCCGCATCGTCTATGGATGTGCCTTTCTGCTTGTTGTCCTGCTGTCCATGCTATGCCTGGTATTGGCCCATTGGCCCAAACGCCTGTTCTCTGGAGAGCTGATCTACATCG
390
Ex1
Ex2
CAT-AmRIA
I P T P R W G R I V Y G C A F L L V V L L S M L C L V L A H W P K R L F S G E L I Y I
ORF-3
I P T P R W G R I V Y G C A F L L V V L L S M L C L V L A H W P K R L F S G E L I Y I
ORF-start-2
SEQ ID NO. 13
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ORF-Start-1
CAGCCGCTGTGTTGCTGCTGGTGCTCATTGTGGGATTCACTTTCACCGTCTGGAGACAGCCGCAGAGCAACACTCCTCTTTACTTCAAGGTCCCCCTGTTGCCTGTTCTCCTCTGGTCAGCATCTCTGCA
520
Ex2
Ex3
CAT-AmRIA
A A A V L L L V L I V G F T F T V W R Q P Q S H T P L Y F K V P L L P V L L W S A S L H
ORF-3
A A A V L L L V L I V G F T F T V W R Q P Q S H T P L Y F K V P L L P V L L W S A S L H
ORF-start-2
S G C V A A G A H C G I H F H R L E T A P E Q H S S L L Q G P P V A C S P L V S I S A
ORF-Start-1

Fig. 7 Part 1

SEQ. ID. No. 10

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CCCAATGTGGAGTCTGGATGTTGATTGGATTTGCTATGTATTTTGGATATGGGATCCCACACAGCTTGGAGAACAATCAA
CAGCCACCAGCGTCAAGCTCCCAAACTCCTCATGAAAACATCCCTAGTCCCTTGCTGAATTAGTTTAACCACGGAAAAGG
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GTTTGTTATAGAGGTGAGGACTTGCTCTGTTGCCCAGGCGGAACTTGAACTCCTGGGCTCAAGTGATCCTCCTGCCTCAG
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Fig. 15

SEQ. ID. No. 13

142 aa

MGPHRLWMCLSACCPAVHAMPGIGPLAQTPVLWRADLHRSGCVAAGAHCGIHFHRLETA
PEQHSSLLQGPPVACSPLVSISAHVYLMMQMTTETWAQCGVWMLIGFAMYFGYGIPHSLE
NNQQPPASSSQTPHENIPSPLLN

Fig. 16

SEQ. ID. No. 14

119 aa

MNSVPKAESPACVPEASSTPASLWSPVSTIPTPRWGRIVYGCAFLLVVLLSMLCLVLAHWP
KRLFSGELIYIAAAVLLLVLIVGFTFTVWRQPQSNTPLYFKVPLLPVLLWSASLHMFT

Fig. 17

SEQ. ID. No. 15

140 aa

FRYQPDENFS KNEKPKEEVV EMNSVPKAES PACVPEASST PASLWSPVST IPTPRWGRIV
YGCAFLLVVL LSMLCLVLAH WPKRLFSGEL IYIAAAVLLL VLIVGFTFTV WRQPQSNTPL
YFKVPLLPVL LWSASLHMFT

Fig. 18

SEQ. ID. No. 11

TGGATATGGGTAGATGAAACCTCGGATACGTGGGAGCTACCGTGGGTGTATTAGGCTC
TCTGATGTTGGATGAACAGGGACCTGGGTTAATCTTGCCCTACTCTCTCTAATTCAGGT
ACCAGCCAGACGAGAATTTCAGCAAGAACGAGAAGCCGAAGGAGGAAGTAGTTGAGA
TGAATTCTGTACCCAAAGCAGAATCCCCAGCATGTGTTCCTGAAGCGTCCAGCACTCCG
GCGAGTCTGTGGAGCCCTGTCAGCACCATCCCCACCCCGAGATGGGGCCGCATCGTCT
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CCAGCGTCAAGCTCCCAAACTCCTCATGAAAACATCCCTAGTCCCTTGCTGAATTAGTT
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TGTGAAGCTCTGTGGAGCATGAAGAGATGGAGTTTCACCTTGTTGCCGAGGCTGGTCTT
GAACTCCTGAGCTGAAGCCATCCTCCTGCCTCTGGCCTCTAAAAGTGCTGGGATTCCAG
GCGTGAGCCAGCGTGCCCGGCCATCATGTTTACTTCAAACAGATTTTCTCTGGCGTCTT
CTAGGACACTTGTATCTGATCTTTACTGAAAGTCACATTCTTAGCAAGACTTAACCTTA
CATCCTGAGTAAAATTGCGAATTCCACAGATGCCCTCAATCTTTCCTGACCTCGGCTCC
TCTTCTGTTTCCCGGCATGGCTTGAAGCGCAACGGCATTCTACCTGTTTACTCAGGATG
TGTCTTACCATTAAAGTCCCTGCCTGCAGAGCAGTTGGCTTGATTTCACCT

Fig. 19

SEQ. ID. No. 12

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GCACTTTGGGAGGCTGAGGCGGGACAACTGCTTGAGCCTGGGCAGTCGAGGCTTCAGTGAATCAAGACCACACCACTGCA
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```

Fig. 20 Part 1

CHARACTERISTIC	Localization/designation
mRNA	Combine (1..312,313..479,480..587,588..1437) /Feature ="CAT-A mRNA Exon 1-4 "
Exon	1..312 /Feature ="CAT-A Exon 1"
Exon	313..479 /Feature ="CAT-A Exon 2"
Exon	480..587 /Feature ="CAT-A Exon 3"
Exon	588..1437 /Feature ="CAT-A Exon 4"
Div_Charakteristikum SEQ ID NO. 13	275..703 /Feature ="CAT-A ORF -Start-1" /Translation="MGPHRLWMCLSACCPAVHAMPGIGPLAQTPVLWRADLHRSGCVAAGAHCGIHFHRLETAPEQHSSLLQGPPVACSPLVSISAHVYLMMQMTTETWAQCGVWMLIGFAMYFGYGIPHSLENNQQPPASSSQTPHENIPSPLLN."
Div_Charakteristikum SEQ ID NO. 14	174..530 /Feature ="CAT-A ORF-start-2" /Translation="MNSVPKAESPACVPEASSTPASLWSPVSTIPTPRWGRIVYGCAFLLVVLLSMLCLVLAHWPKRLFSGELIYIAAAVLLLVLIVGFTFTVWRQPQSNTPLYFKVPLLPVLLWSASLHMFT"
SEQ ID NO. 15	Div_Charakteristikum 111..530 /Feature ="CAT-A ORF-3" /Translation="FRYQPDENFSKNEKPKEEVVEMNSVPKAESPACVPEASSTPASLWSPVSTIPTPRWGRIVYGCAFLLVVLLSMLCLVLAHWPKRLFSGELIYIAAAVLLLVLIVGFTFTVWRQPQSNTPLYFKVPLLPVLLWSASLHMFT"
Div_Charakteristikum	482..667 /Feature ="58% homology to humans -CAT3 GB Acc. AF320612"
Div_Charakteristikum	482..667 /Feature ="63% Protein homology to cationic amino acid transporter 3 [Rattus norvegicus] GB Acc.BAA20133.1"
Div_Charakteristikum	452..682 /Feature ="protein homology to ecotropic retrovirus receptor [Rattus norvegicus] GB Acc. BAB83893"
Div_Charakteristikum	24..500 /Feature ="hom-rat-baa20133.1-31%"
Div_Charakteristikum	24..500 /Feature ="hom-mouse-cat3-AAB09593-60%"
Div_Charakteristikum	482..667 /Feature ="hom-mouse-cat3-AAB09593-40%"
Div_Charakteristikum	500..682 /Feature ="hom mouse-cat2-AAA37372-42%"

NUMBER OF BASES 333 a 345 c 341 g 415 t

SEQ ID NO. 10

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GCACTCCGGCGAGTCTGTGGAGCCCTGTCAGCACCATCCCCACCCCGAGATGGGGCCGCATCGTCTATGGATGTG
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TTACCTGATGATGCAGATGACCACTGAGACCTGGGCCCAATGTGGAGTCTGGATGTTGATTGGATTTGCTATGTA
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GCAAACACTTTTTAAAAAACTTTATTTTGTTGTTTGTTTGTTTGTTTGTTTGTTTGTTTGTTATAGAGGTGAGGA
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Fig. 21 Part 1

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CACCACCACGCCCGGCTAATTTTTGTATTTTTAGTAGAGATGGGGTTTCAACATGGTGGCCAGGCTGGTCTGGAA
CTCCTGACCTCGTGATCTGCCCACCTTGGCCTCCCAAAGTGCTGGGACTACAGAAGTGAGCCACCGCGCCTGGCT
TGTATTTTCTTAAAGAAATAAAAAACTAAGAAGAAAATGCCACTCATCATCATTGGTATAGCAATGTGGTCAGCA
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GGTTTTTCACTAGATATTTCTTTTACTGCTGCTCTTCCTTCTTTTCTACATCTCATATTAGAGGATTGGACAGCC
TTTAAAACTTCATTGACAGGAGCAAATCCAGTATGCACTGATTTCTTTTTTTTTTTCTTTTTTTTTTTTTTGGTG
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TAGGCCAAGAAAGCTACTGGCCTCTGATGCCCTTGTTCATCAGTCATAGAGCCAACACCTGGCGGTTCAACCGTA
ACATCATAAACGATTCCTCTGGTGATGAGGAAGTAAGACACCACCACCAGAGCATACATAGTCATGGCCAACGGC
ATGTGCAGCCAGGGCGGCTTCTTCAGCTTCAGGTTGGGACATTCAAGCGCTAAGAACGGGACACGGTACAAAGTC
TCCATGTTGGTGGCAGCAAGGGCCATTCTCGGCCTC

SEQ ID NO. 16

Fig. 22

CHARACTERISTIC

Exon	Localization/designation 1..288
Div_Charakeristikum	/Feature ="Exon 3" complementary (1..288) /Feature =" homologous to XM_034710-mRNA-782-1070 end section.seq(1>288)"
Exon	621..863
Div_Charakeristikum	/Feature ="exon 2" complementary (621..863) /Feature =" homologous to XM_034710-mRNA-539-782-part.seq(1>244)"
Exon	1183..1686
Div_Charakteristikum	/Feature ="Exon 1" complementary (1183..1686) /Feature =" homologous to XM_034710-mRNA-37-538-part.seq(1

NUMBER OF BASES 426 a 403 c 350 g 507 t

TTTTTTAATTAGTATTTATTTCTGTTCCTTTCATACATGTTATTGTTCAGGCAGCAGTCTTACATACCTATCTGA
AATAATCTACCATCACACACTCTAGTTAAAGCCAAAGCTCCGTAGGTAAAGCTGACCAGCTGTCCAGAGTTCTCA
GCTCATGGAATCTTGCTTCTTGATTTTATGTTAATTTCTGAAAGTCAGAAATGTCATCTCCAAAAAGTGTTAAGG
GGGTTGTAAAAAATAATCACTATATTTGCTATTATAGTTGTAGGTATGCAGTATATTTTCTTTTCTTTCTTTCTT
TCTTTTTTTTTTTTTTGAAACAGAGTCTCGCTCTGTCGCCCAGGCTGGAGTGCAGTGGCAGCGATCTAGGCTCACT
GCAGCCTCCGCCTCCCAGGTTCACGTGATTCTCCTGCCTCAGCCTTCCAAGTAGCTGGGATTACAGGTGCCCCGC
CACCACCACGCCCGGCTAATTTTTGTATTTTTAGTAGAGATGGGGTTTCAACATGGTGGCCAGGCTGGTCTGGAA
CTCCTGACCTCGTGATCTGCCCACCTTGGCCTCCCAAAGTGCTGGGACTACAGAAGTGAGCCACCGCGCCTGGCT
TGTATTTTCTTAAAGAAATAAAAAACTAAGAAGAAAATGCCACTCATCATCATTGGTATAGCAATGTGGTCAGCA
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SEQ ID NO. 16

Fig. 23

CHARACTERISTIC	Localization/designation
Exon	748..1035 /Feature ="Exon 3"
Div_Charakteristikum	748..1035 /Feature =" homologous to XM_034710-mRNA-782-1070-end section.seq(1>288)"
Exon	505..747 /Feature ="Exon 2"
Div_Charakteristikum	505..747 /Feature =" homologous to XM_034710-mRNA-539-782-part.seq(1>244)"
Div_Charakteristikum	1..504 /Feature ="Exon 1"
Div_Charakteristikum	1..504 /Feature =" homologous to XM_034710-mRNA-37-538-part.seq(1>502)"

NUMBER OF BASES 307 a 194 c 222 g 312 t

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CAACCTGAAGCTGAAGAAGCCGCCCTGGCTGCACATGCCGTTGGCCATGACTATGTATGCTCTGGTGGTGGTGTC
TTACTTCCTCATCACCAGAGGAATCGTTTATGATGTTACGGTTGAACCGCCAGGTGTTGGCTCTATGACTGATGA
ACAAGGGCATCAGAGGCCAGTAGCTTTCTTGGCCTACAGAGTAAGTGGACAATATTATTATGGAAGGACTTGGAT
CCAGCTTCCTGTTTACAATGGGAGGTTTAGGTTTCATAATCCTGGACCGATCGAATGCACCAAATATCCCAAAAC
TCAATAGATTTCTTCTTCTATTCGTTGGATTCGTCTGTGTCCTATTGAGTTTTTTTCACGGCTAGAGTATTCGTG
AGAATGAAACTACCGGGCTCTCTGATGGGTTAGAGTGCCTTTAAGAAGAAATCAGTGCATACTGGATTTGCTCCT
GTCAATGAAGTTTTAAAGGCTGTCCAATCCTCTAATATGAGATGTAGAAAAGAAGGAAGAGCAGCAGTAAAAGAA
ATATCTAGTGAAAAACCAGGAAGTGTATTGAAGCTTGGACTAGAATTTCTTCTTTATTAAAGAGACAAATTTATC
ACAGTATTTTCTTTTCCTGCTGACCACATTGCTATACCAATGATGATGAGTGGCATTTTCTTCTTAGTTTTTAAA
GAAAATATACTGCATACCTACAACTATAATAGCAAATATAGTGATTATTTTTTACAACCCCCTTAACACTTTTTG
GAGATGACATTTCTGACTTTCAGAAATTAACATAAAATCAAGAAGCAAGATTCCATGAGCTGAGAACTCTGGACA
GCTGGTCAGCTTTACCTACGGAGCTTTGGCTTTAACTAGAGTGTGTGATGGTAGATTATTTCAGATAGGTATGTA
AGACTGCTGCCTGAACAATAACATGTATGAAAGGAACAGAAATAAATACTAATTAAAAAA
```

SEQ ID NO. 17

Fig. 24

CHARACTERISTIC	Localization/designation
Exon	2118..2524 /Feature ="Exon 18"
Exon	1976..2117 /Feature ="Exon 17 "
Exon	1867..1975 /Feature ="Exon 16 "
Exon	1785..1866 /Feature ="Exon 15 "
Exon	1645..1784 /Feature ="Exon 14 "
Exon	1581..1644 /Feature ="Exon 13 "
Exon	1488..1580 /Feature ="Exon 12 "
Exon	1298..1487 /Feature ="Exon 11 "
Exon	1144..1297 /Feature ="Exon 10"
Exon	1113..1143 /Feature ="Exon 9"
Exon	1024..1112 /Feature ="Exon 8"
Exon	888..1023 /Feature ="Exon 7"
Exon	730..887 /Feature ="Exon 6"
Exon	597..729 /Feature ="Exon 5"
Exon	484..596 /Feature ="Exon 4"
Exon	400..483 /Feature ="Exon 3"
Exon	367..399 /Feature ="Exon 2"
Exon	1..366 /Feature ="Exon 1"

NUMBER OF BASES	475 a	814 c	729 g	506 t

SEQ ID NO. 19

```
GGCGCGCCCGGGGTGCCGCTCCCTGCCTGGCGCGCTCCGCACCTGGAGGTGCCTTGCCCCTCTCCTGCCCACCTC
GGAATTTCCCTGTGGCTCCTTTGATCCTTCGAGTCTCCAGCTCCTCTCCCTTCCACCTGTTTCCCCCAAGAAAGG
CAGGATCCTGGTCCCTGCTACGTTTCTGGGGCCATGGCTGGTCTGGGCCCCGGCGTAGGCGATTCAGAGGGGGGA
CCCCGGCCCCTGTTTTGCAGAAAGGGGGCCCTGAGGCAGAAGGTGGTCCACGAAGTCAAGAGCCACAAGTTCACC
GCTCGCTTCTTCAAGCAGCCCACCTTCTGCAGCCACTGCACCGACTTCATCTGGTGAGGGAAGGGGGCAGGGCCT
GCAATGTCAAGGTAAGAGCTGGGGTGGTTCATCGACGATGCCACGAATTTGTGACCTTCGAGTGTCCAGGCGCTG
GGAAGGGCCCCCAGACGGACGTGAGTGCTCGGACCGGAACAAACACAAGTTCCGCCTGCATAGCTACAGCAGCCC
CACCTTCTGCGACCACTGTGGCTCCCTCCTCTACGGGCTTGTGCACCAGGGCATGAAATGCTCCTGTGAGTGAGA
TGAACGTGCACCGGCGCTGTGTGCGTAGCGTGCCCTCCCTGTGCGGTGTGGACCACACCGAGCGCCGCGGGCGCC
TGCAGCTGGAGATCCGGGCTCCCACAGCAGATGAGATCCACGTAACTGGTGAGGGAGGCCCGTAACCTAATTCCT
ATGGACCCCAATGGTCTCTCTGATCCCTATGTGAAACTGAAGCTCATCCCAGACCCTCGGAACCTGACGAAACAG
AAGACCCGAACGGTGAAAGCCACGCTAAACCCTGTGTGGAATGAGACCTTTGTGTTGTGAGTCCTGAAGCCAGGG
GATGTGGAGCGCCGGCTCAGCGTGGAGGTGTGGGACTGGGACCGGACCTCCCGCAACGACTTCATGGGGGCCATG
TCCTTTGGCGTCTCGGAGCTGCTCAAGGCGCCCGTGGATGGCTGGTGACAAGTTACTGAACCAGGAGGAGGGCGA
GTATTACAATGTGCCGGTGGCCGATGCTGACAACTGCAGCCTCCTCCAGAAGTTTGAGGTACTGTAACTACCCCC
TGGAATTGTATGAGGTGAGCGGGTGCGGATGGGCCCCTCTTCCTCTCCCATCCCCTCCCCTTCCCCTAGTCCCAC
CGACCCCAAGCGCTGCTTCTTCGGGGCGAGTCCAGGACGCCTGCACATCTCCGACTTCAGCTTCCTCATGGTTCT
AGGAAAAGGCAGTTTTGGGAAGGGTGATGCTGGCCGAGCGCAGGGGCTCTGATGAGCTCTACGCCATCAAGATCT
TGAAAAAGGACGTGATCGTCCAGGACGACGATGTGGACTGCACGCTGGTGGAGAAACGTGTGCTGGCGCTGGGGG
GCCGGGGTCCTGGCGGCCGGCCCCACTTCCTCACCCAGCTCCACTCCACCTTCCAGACCCCGGACCGCCTGTATT
TCGTGATGGAGTACGTCACCGGGGGAGACTTGATGTACCACATTCAACAGCTGGGCAAGTTTAAGGAGCCCCATG
CAGCGGTTCTACGCGGCAGAAATCGCTATCGGCCTCTTCTTCCTTCACAATCAGGGCATCATCTACAGGGGACCT
GAAGCTGGACAATGTGATGCTGGATGCTGAGGGACACATCAAGATCACTGACTTTGGCATGTGTAAGGAGAACGT
CTTCCCCGGGACGACAACCCGCACCTTCTGCGGGACCCCGGACTACATAGCCCCGGAGGATCATTGCCTACCAGC
```

Fig. 25 Part 1

CHARACTERISTIC	Localization/designation
Source	complementary (1..24801) /Organismus="Homo sapiens" /db_xref="taxon:9606" /Chromosome="19" /Clone="CTC-331H23"
Exon	24395..24801 /Feature ="Exon 18 "
Exon	24005..24146 /Feature ="Exon 17"
Exon	22324..22432 /Feature ="Exon 16"
Exon	20762..20843 /Feature ="Exon 15"
Exon	18300..18439 /Feature ="Exon 14"
Exon	18108..18171 /Feature ="Exon 13"
Exon	17922..18014 /Feature ="Exon 12"
Exon	16128..16317 /Feature ="Exon 11"
Exon	15647..15800 /Feature ="Exon 10"
Exon	11054..11084 /Feature ="Exon 9"
Exon	10680..10768 /Feature ="Exon 8"
Exon	10201..10336 /Feature ="Exon 7"
Exon	9368..9525 /Feature ="Exon 6"
Exon	7580..7712 /Feature ="Exon 5"
Exon	7332..7444 /Feature ="Exon 4"
Exon	1862..1945 /Feature ="Exon 3 "
Exon	864..896 /Feature ="Exon 2"
Exon	1..366 /Feature ="Exon 1"

NUMBER OF BASES 5613 a 6407 c 6621 g 6160 t

SEQ ID NO. 18

```
GGCGCGCCCGGGGTGCCGCTCCCTGCCTGGCGCGCTCCGCACCTGGAGGTGCCTTGCCCCTCTCCTGCCCACCTC
GGAATTTCCCTGTGGCTCCTTTGATCCTTCGAGTCTCCAGCTCCTCTCCCTTCCACCTGTTTCCCCCAAGAAAGG
CAGGATCCTGGTCCCTGCTACGTTTCTGGGGCCATGGCTGGTCTGGGCCCCGGCGTAGGCGATTCAGAGGGGGGA
CCCCGGCCCCTGTTTTGCAGAAAGGGGGCCCTGAGGCAGAAGGTGGTCCACGAAGTCAAGAGCCACAAGTTCACC
GCTCGCTTCTTCAAGCAGCCCACCTTCTGCAGCCACTGCACCGACTTCATCTGGTGAGGGAAGGGGGCTGGGGGA
CTGGGGGACGAGGGGACTAGGGGTGCAGACTCCTATCACGCCGACCCCTGTGGAAGGAAGAAGGAGGGGGCTGTA
GTCCCGACTCCCAGGTTCTAGGATGGCCAGGGAACGCTGGGAGCTTCGACTCCTGGGTTTCAGTGAGGAGGAGGC
TGGTTCCTGGAGTGCTGGGTCCGAGGGAGGAGGAGGCTGGAGGACAGAGGTCCTGGAGTCTTGGGTCTGAGGGAG
GAAGGGCCTGGGGGGCTGGGAGCCTGGATTCCTGGGTCTGAAGGAGGAAGAAACTGGGGGCTGAACTCCAGTCTA
AGGGAAGAAGGGCTGGGGGCCAAAATTTCTGGGTTCTAGAAAGAGGAGGTGGCCGGGGCTTGGACACCTGGGCCC
TGCGGGAGGAGGGTCAGAGAGCGCAGGCCCCCTGTGGCTCGCAGAGGTTGGGGGTCCAGGTACCCCTTTCTGCAC
TGACCTAGGATCCCTGACTCTTCCAGGGGTATCGGAAAGCAGGGCCTGCAATGTCAAGGTAAGAGCTGGGGACCG
GGGCTCCTGGGACCCTCAGGAGGGTGGAGGCTGGGGCCCCACAGCTGAGGCTGCTTGACACACGTGTTCTCTGGT
CCCCAGAGAGGCGCGGGGGAGCCCGGGGCGGGGGGTGTGGCAGAGACACAGCCTGTGGTGGGGAGGGAGCTTTGA
TGGTGGGGCCACCGCGGAGGTGGTGCTGGGGGCCCCTCCCTCGGCCGGCTCCAGGTGGGGACAGATATTTGGAGA
ATCTGTTGCCATGGGAACATGGAGATTTGGAAAAGGGGAGCTGAAGGGGGGAAGGGGGAGGGGGCTGGAGATGCA
AAGTCAGAGCCCCCCCCCACCCCAGGCTGCCGTCGCCATGACAACACCAGCCGTCCTAGGCAGGGGCAGGCCGGG
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Fig. 26 Part 1